

## ENTERED

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/905,743B

DATE: 04/21/2003 TIME: 13:14:11

Input Set : N:\EBONY'S\EP.txt

Output Set: N:\CRF4\04212003\I905743B.raw

3 <110> APPLICANT: Chadwick, Brian Paul Frischauf, Anna Maria 6 <120> TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO CD39-LIKE POLYPEPTIDES AND NUCLEIC ACIDS 9 <130> FILE REFERENCE: 28110/36120C 11 <140> CURRENT APPLICATION NUMBER: 09/905,743B 12 <141> CURRENT FILING DATE: 2001-07-13 14 <150> PRIOR APPLICATION NUMBER: 09/240,639 15 <151> PRIOR FILING DATE: 1999-01-29 17 <160> NUMBER OF SEQ ID NOS: 32 19 <170> SOFTWARE: PatentIn version 3.1 21 <210> SEQ ID NO: 1 22 <211> LENGTH: 2762 23 <212> TYPE: DNA 24 <213> ORGANISM: Homo Sapiens 26 <220> FEATURE: 27 <221> NAME/KEY: CDS 28 <222> LOCATION: (232)..(1599) 29 <223> OTHER INFORMATION: W--> 32 < 400 > 1\_33 gtggggtegt atecegeggg tggaggeegg ggtggegeeg geeggggegg gggageeeaa 60 35 aagaccggct geegeetget eeeeggaaaa gggeaetegt eteegtgggt gtggeggage 120 37 gcgcggtgca tggaatgggc tatgtgaatg aaaaaaggta tccgttatga aacttccaga 180 39 aaaacgaget acatttttca geageegeag caeggteett ggeaaacaag g atg aga 237 40 Met Arg 41 285 43 aaa ata too aac cac ggg agc otg ogg gtg gog aag gtg gca tac ooc 44 Lys Ile Ser Asn His Gly Ser Leu Arg Val Ala Lys Val Ala Tyr Pro 10 333 47 ctg ggg ctg tgt gtg ggc gtg ttc atc tat gtt gcc tac atc aag tgg 48 Leu Gly Leu Cys Val Gly Val Phe Ile Tyr Val Ala Tyr Ile Lys Trp 51 cac cgg gec acc gec acc cag gec tte tte age ate acc agg gea gec 381 52 His Arg Ala Thr Ala Thr Gln Ala Phe Phe Ser Ile Thr Arg Ala Ala 53 35 40 50 55 ccg ggg gcc cgg tgg ggt cag cag gcc cac agc ccc ctg ggg aca gct 56 Pro Gly Ala Arg Trp Gly Gln Gln Ala His Ser Pro Leu Gly Thr Ala 55 60 59 gca gac ggg cac gag gtc ttc tac ggg atc atg ttt gat gca gga agc 477 60 Ala Asp Gly His Glu Val Phe Tyr Gly Ile Met Phe Asp Ala Gly Ser 75 525 63 act ggc acc cga gta cac gtc ttc cag ttc acc cgg ccc ccc aga gaa

64 Thr Gly Thr Arg Val His Val Phe Gln Phe Thr Arg Pro Pro Arg Glu

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69	100					105					110					
71 tct																621
72 Ser	Ala	Tyr	Ala	Asp	Asp	Val	Glu	Lys	Ser	Ala	Gln	Gly	Ile	Arg	Glu	
73 115					120					125					130	
75 cta	_	-	_	_		_	_		_		-			_	_	669
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77				135					140					145		
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99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr	y Sei c tco a Sei c aao	Thr c cca r Pro 245 g cto	Gln 230 ccc Pro tat	atc Ile ggc Gly	tac Tyr	ctg Leu agc	acg Thr 250 tac	Pro 235 gca Ala	cgc Arg ctg Leu	Val cgg Arg	Glu gatg Met ggg Gly	ttt The 255 ctg	Thi 240 aac Asr atg	ctg Leu agg Arg	Gln acc Thr	1005
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109	y Ser c too a Ser c aao c Lys 260	Thr c cca r Pro 245 g cto s Leu	Gln 230 ccc Pro tat	atc Ile ggc Gly tcc	tac Tyr tac	ctg Leu agc Ser 265	acg Thr 250 tac	Pro 235 gca Ala cto	cgc Arg Ctg Ctg Ctg Ctg Ctg	Val g cgg g Arg g ctc g Leu	Glumatom Met Gly Gly Gly 270	ttt The 255 ctg	Thi 240 aao Asr ato Met	ctg Leu cagg Arg tcg	Gln acc Thr gca Ala	1005 1053
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109 111 cgc	Y Ser c too ser c aao c Lys 260 c cto	r Thr c cca r Pro 245 g cto s Leu )	Gln 230 ccc Pro tat Tyr	atc Ile ggc Gly tcc Ser	tac Tyr tac Tyr	ctg Leu agc Ser 265	acg Thr 250 tac Tyr	Pro 235 gca Ala cto Leu	cgc Arg ctg Ctg Ctg Ctg Gly	y Val y cgg n Arg y ctc y Leu	atg Met ggg ggg ggg 270	ttt Phe 255 ctg Leu	Thi 240 aac Asr ato Met	ctg Leu agg Arg tcg Ser	Gln acc Thr gca Ala gga	1005
99 gga 100 Gly 101 103 gcd 104 Ala 105 107 tag 108 Tyr 109 111 cgg 112 Arg	y Sen c too Sen c aag c Lys 260 c cto	r Thr c cca r Pro 245 g cto s Leu )	Gln 230 ccc Pro tat Tyr	atc Ile ggc Gly tcc Ser	tac Tyr tac Tyr	ctg Leu agc Ser 265 ggc	acg Thr 250 tac Tyr	Pro 235 gca Ala cto Leu	cgc Arg ctg Ctg Ctg Ctg Gly	y Val y cgg y Arg y ctc y Leu y cag	atgrater Met Gly Met Gly Gly 270	ttt Phe 255 ctg Leu	Thi 240 aac Asr ato Met	ctg Leu agg Arg tcg Ser	Gln acc Thr gca Ala gga Gly	1005 1053
99 gga 100 Gly 101 103 gcd 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275	y Ser c too a Ser c aao c Lys 260 c cto	Three coarse Process Leur	GIn 230 ccc Pro tat Tyr atc	atc Ile ggc Gly tcc Ser	tac Tyr ggc Gly 280	ctg Leu agc Ser 265 ggc Gly	acg Thr 250 tac Tyr	Pro 235 gca Ala cto Leu gag	cgc Arg ctg Leu ggg Gly	y Val y cgg y Arg y ctc y Leu y cag y Gln 285	atg Met Gly 270 cct	ttt Phe 255 ctg Leu gct	Thr 240 aac Asr datg Met	ctg Leu cagg Arg tcg Ser gat Asp	Gln acc Thr gca Ala gga Gly 290	1005 1053 1101
99 gga 100 Gly 101 103 gcd 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac	y Ser c too a Ser c aag c Lys 260 c cto tec too g Lec	Three ccare Process Leury gcg I Ala	GIN 230 CCC Pro CCC Tark Tyr atc	atc Ile ggc Gly tcc Ser ctg Leu agc	tac Tyr tac Tyr ggc Gly 280	ctg Leu agc Ser 265 ggc Gly	tace Tyr  gtg Val	Pro 235 gca Ala cto Leu gag Glu	cgc Arg	y Val y cgg i Arg y ctc y Leu y cag y Gln 285 c agt	g atg y atg y Met g ggg 270 y cct pro	ttt Phe 255 ctg Leu gct Ala	Thi 240 aac Asr ato Met	ctg Leu cagg Arg tcg Ser sat gat sat	Gln acc Thr gca Ala gga Gly 290 tgg	1005 1053
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac 116 Lys	y Service took Service Lys 260 ctook	Three ccar pro 245 cto s Leu ) gcg la Ala	Gln 230 ccc Pro Ctat Tyr atc Ile	atc Ile ggc Gly tcc Ser ctg Leu agc	tac Tyr tac Tyr ggc Gly 280 cct Pro	ctg Leu agc Ser 265 ggc Gly tgc	acg Thr 250 tac Tyr gtg Val	235 gca Ala cto Leu gag Glu tct	cgc Arg Ctg Ctg Ctg Gly Gly Ccc Pro	y Val y cgg y Arg y ctc y Leu y cag y Gln 285 e agt	g atg y atg y Met g ggg 270 270 cct h Pro	ttt Phe 255 ctg Leu gct Ala	Thi 240 aac Asr ato Met	ctg Leu ctg Leu cagg Arg cs Ser gat sAsp	Gln acc Thr gca Ala Gly 290 tgg Trp	1005 1053 1101
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac 116 Lys	y Ser c too c aag c Lys 260 c cto g Leo g gao g Glu	Three coare Process Leury goog good all Ala	GIN 230 CCC Pro CCC Tatal Tyr atc Ile	atc Ile ggc Gly tcc Ser ctg Leu agc Ser 295	tac Tyr tac Tyr ggc Gly 280 cct	ctg Leu agc Ser 265 ggc Gly tgc	acg Thr 250 tac Tyr gtg Val	Pro 235 gca Ala cto Leu gag Glu tct	cgc Arg Ctg Ctg Ctg Ctg Cgc Cgc Cgc Cgc Cgc Cgc Cgc Cgc Cgc Cg	y Val y cgg y Arg y ctc y Leu y cag y Gln 285 c agt	g atg met g ggg g Gly 270 g cct pro	g ttt Phe 255 g ctg Leu gct Ala aaa Lys	Thr 240 aac Asr atg Met Laag Lys	ctg Leu cagg Arg stcg Ser gat Asp Glu 305	Gln acc Thr gca Ala gga Gly 290 tgg Trp	1005 1053 1101 1149
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac 116 Lys 117	y Ser c too c aag c Lys 260 c cto g Leu 5 g gao g Glu	Three coarse Proceedings Level 19 19 19 19 19 19 19 19 19 19 19 19 19	Gln 230 ccc Pro Ctat Tyr atc Ile Val	atc Ile ggc Gly tcc Ser ctg Leu agc Ser 295 gtc	tac Tyr tac Tyr ggc Gly 280 cct Pro	ctg Leu agc Ser 265 ggc Gly tgc Cys	acg Thr 250 tac Tyr gtg Val ttg Leu	Pro 235 gca Ala cto Leu gag Glu tct Ser	cgc Arg	y Val y cgg y ctc y Leu y cag y Gln 285 c agt	atg Met ggg Gly 270 cct Pro	To Gly The	Thr 240 aac Asr atg Met Lys gga Gly	ctg Leu cagg Arg stcg Ser gat agag Glu 305	Gln acc Thr gca Ala gga Gly 290 tgg Trp gca	1005 1053 1101
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac 116 Lys 117 119 gaa 120 Glu	y Ser c too c aag c Lys 260 c cto g Leu 5 g gao g Glu	Three coarse Proceedings Level 19 19 19 19 19 19 19 19 19 19 19 19 19	Gln 230 ccc Pro ctat Tyr atc Ile Val	atc Ile ggc Gly tcc Ser ctg Leu agc Ser 295 gtc Val	tac Tyr tac Tyr ggc Gly 280 cct Pro	ctg Leu agc Ser 265 ggc Gly tgc Cys	acg Thr 250 tac Tyr gtg Val ttg Leu	Pro 235 gca Ala cto Leu gag Glu tct Ser	cgc Arg	y Val y cgg y ctc y Leu y cag y Gln 285 c agt	atg Met ggg Gly 270 cct Pro	To Gly The	Three 240 and a Asri Met Lys Gly Gly Gly Gly Alas	ctg Leu cagg Arg scar gag scar gag scar gag scar gag scar gag scar agg scar	Gln acc Thr gca Ala gga Gly 290 tgg Trp gca	1005 1053 1101 1149
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac 116 Lys 117 119 gaa 120 Glu	y Service too too too too too too too too too to	Three coars are produced as Leuchard Alace goals Alace	Gln 230 ccc Pro tat Tyr atc Ile gtc Val gaa Glu 310	atc Ile ggc Gly tcc Ser ctg Leu agc Ser 295 gtc Val	tac Tyr tac Tyr ggc Gly 280 cct Pro	ctg Leu agc Ser 265 ggc Gly tgc Cys	acg Thr 250 tac Tyr Val ttg Leu	Pro 235 gca Ala cto Leu gag Glu tct Ser gtt Val 315	cgc Arg	y Val y cgg y ctc y Leu y cag y Gln 285 agt Ser	g atg Met g ggg Gly 270 cct Pro	THE COLUMN	Thir 240 aac Asr at a Met Lys Gly	ctg Leu cagg Arg scar gag scar gag scar gag scar gag scar agg scar	Gln acc Thr gca Ala gga Gly 290 tgg Trp gca Ala	1005 1053 1101 1149
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac 116 Lys 117 119 gaa 120 Glu	y Service too too too too too too too too too to	Three coars are process Leuron process Leuron process Leuron process Leuron process Leuron process Leuron process Alaman proce	Gln 230 ccc Pro tat Tyr atc Ile gtc Val gaa Glu 310 gag	atc Ile ggc Gly tcc Ser ctg Leu agc Ser 295 gtc Val	tac Tyr tac Tyr ggc Gly 280 cct Pro	ctg Leu agc Ser 265 ggc Gly tgc Cys tac Tyr	acg Thr 250 tac Tyr Val ttg Val agg	Pro 235 gca Ala cto Leu gag Glu tct Ser yal 315 aga	cgc Arg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ct	y val y cgg y ctc y Leu y cag y Gln 285 agt y agt y ggg	gaggagagagagagagagagagagagagagagagagag	To Gly The State of Ala The Al	Thir 240 and a control of the contro	ctg ctg ctg cleu cagg cagg cagg cagg cagg cagg cagg cag	Gln acc Thr gca Ala gga Gly 290 tgg Trp gca Ala aac	1005 1053 1101 1149
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac 116 Lys 117 119 gaa 120 Glu	y Service too too too too too too too too too to	Three coars are process Leuron process Leuron process Leuron process Leuron process Leuron process Leuron process Alaman proce	Gln 230 Pro Fro Tark I I I I I I I I I I I I I I I I I I I	atc Ile ggc Gly tcc Ser ctg Leu agc Ser 295 gtc Val	tac Tyr tac Tyr ggc Gly 280 cct Pro	ctg Leu agc Ser 265 ggc Gly tgc Cys tac Tyr	acg Thr 250 tac Tyr Val ttg Val agg	Pro 235 gca Ala cto Leu gag Glu tct Ser ytt Val 315 aga Arg	cgc Arg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ct	y val y cgg y ctc y Leu y cag y Gln 285 agt y agt y ggg	gaggagagagagagagagagagagagagagagagagag	To Gly The State of Ala The Al	Thir 240 aac Asr at a Met Lys a gga a Gly a Lys a Lys a gca a 320 ctt. Leu	ctg ctg ctg cleu cagg cagg cagg cagg cagg cagg cagg cag	Gln acc Thr gca Ala gga Gly 290 tgg Trp gca Ala aac	1005 1053 1101 1149
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac 116 Lys 117 119 gaa 120 Glu 121 123 agc 124 Ser 125	y Server took took took took took took took too	Three coactions are the coactions and the coactions are the coacti	Gln 230 CCC Pro CCC Tata Tyr atc Ile Glu 310 Glu 310 Glu	atc Ile ggc Gly tcc Ser ctg Leu agc Ser 295 gtc Val	tac Tyr tac Tyr ggc Gly 280 cct Pro acg Thr	ctg Leu agc Ser 265 ggc Gly tgc Cys tac Tyr	acg Thr 250 tac Tyr Val ttg Val agg Arg	gag Glu tct Ser gtt Val 315 aga Arg	cgc Arg Ctg Ctg Ctg Ctg Cgc Cgc Cgc Cgc Cgc Cgc Cgc Cgc Cgc Cg	y Val y Cgg y Arg y Ctc y Leu y Cag y Gln 285 x agt y Ser y Gly y Cgg y Cgly y Cgg y Cgly y Cgg y Cgly y Cgg y y Cgg y Cgg y Cgg y Cgg y y Cgg y y Cgg y y Cgg y y Cgg y	gagger Glu	Total Gly The State Control Th	Three 240 and a control of the contr	ctg ctg ctg cagg cagg cagg cagg cagag caga	Gln acc Thr gca Ala gga Gly 290 tgg Trp gca Ala aac Asn	1005 1053 1101 1149
99 gga 100 Gly 101 103 gcc 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac 116 Lys 117 119 gaa 120 Glu 121 123 agc 124 Ser	y Server took took took took took took took too	Three coactions are the coactions and the coactions are the coacti	Gln 230 ccc Pro Tyr atc Ile Glu 310 gag Glu 320 cc agg	atc Ile ggc Gly tcc Ser ctg Leu agc Ser 295 gtc Val ctg Leu acg	tac Tyr tac Tyr ggc Gly 280 cct Pro acg Thr	ctg Leu agc Ser 265 ggc Gly tgc Cys tac Tyr gct Ala	acg Thr 250 tac Tyr Val ttg Val agg Arg	Pro 235 gca Ala cto Leu gag Glu tct Ser ytt Val 315 aga Arg	cgc Arg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ct	y Val y Cgg y Arg y Ctc y Leu y Cag y Gln 285 c agt y Ser y Gly t Ca y Ser y Gly	gagger Gluman ga	To Gly The State Lys The State	Three 240 and a control of the contr	ctg ctg ctg cleu cagg cagg cagg cagg cagg cagg cagg cag	Gln acc Thr gca Ala gga Gly 290 tgg Trp gca Ala aac Asn	1005 1053 1101 1149 1197
99 gga 100 Gly 101 103 gcd 104 Ala 105 107 tac 108 Tyr 109 111 cgc 112 Arc 113 275 115 aac 116 Lys 117 119 gaa 120 Glu 121 123 agc 124 Ser 125 127 aga	y Server took took took took took took took too	Three coace Process Leur Process Leur Process Leur Process Leur Process Leur Process Alace Process A	Gln 230 ccc Pro Tyr atc Ile Glu 310 gag Glu 320 cc agg	atc Ile ggc Gly tcc Ser ctg Leu agc Ser 295 gtc Val ctg Leu acg	tac Tyr tac Tyr ggc Gly 280 cct Pro acg Thr	ctg Leu agc Ser 265 ggc Gly tgc Cys tac Tyr gct Ala	acg Thr 250 tac Tyr gtg Val ttg Leu agg Arg	Pro 235 gca Ala cto Leu gag Glu tct Ser ytt Val 315 aga Arg	cgc Arg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ctg Ct	y Val y Cgg y Arg y Ctc y Leu y Cag y Gln 285 c agt y Ser y Gly t Ca y Ser y Gly	gagger Gluman ga	To Gly The Phe 255 The Control of Ala The Ala	Three 240 and a control of the contr	ctg ctg ctg cleu cagg cagg cagg cagg cagg cagg cagg cag	Gln acc Thr gca Ala gga Gly 290 tgg Trp gca Ala aac Asn	1005 1053 1101 1149 1197

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	Lys Gly Gly Ser Leu Val Val Gly Asp Phe Glu Ile Ala Ala Lys Tyr	
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210	20 25 30	

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218		50					55			_		60			_	
		Ala	Ala	Asp	GLy		GLu	Val	Phe	Tyr	_	He	Met	Phe	Asp	
222		_	_,	~ 7	_,	70	1	•			75	-1	m1	_	_	80
	Gly	Ser	Thr	GLY		Arg	Val	His	Val		GIn	Phe	Thr	Arg	Pro	Pro
226	_	~ 1	1	_	85	_			<b>~</b> 1	90	<b>m</b> 1.		. 1 .	7	95	
	Arg	Glu	Thr		Thr	Leu	Thr	HIS		Thr	Pne	гàг	Ата		Lys	Pro
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234	3	<b>~1</b>	115	T	7	17.0 ]	<b>31</b> -	120	<b>a</b> 15	7 00	т1.	Dwo	125	7 an	Dho	m~~
	Arg	130	ьeu	ьeu	ASP	Val	135	гуѕ	GIII	ASp	тте	140	Pne	ASP	Phe	пр
238	T ***		mb∞	Dro	Tou	Wa I		T 110	λla	Пhr	λla		LOU	7 ~~	Leu	Lou
	145	ніа	1111	PIO	Leu	150	ьец	цуъ	нта	1 111	155	GLY	Leu	AIY	ьeu	160
		Clv	Glu	Luc	λla		Luc	T.OII	T.011	Gln.		Va 1	Lve	Glu	Val	
246	PIO	GLY	GIU	цуз	165	GIII	цуз	пец	шеu	170	цуз	Vai	шуз	GIU	175	rne
	T.vc	Δla	Ser	Pro		T.011	Va 1	Glv	Δsn		Cvs	Va 1	Ser	Tle	Met	Δsn
250	- Нуз		DCI	180	1110	пси	vai	O <sub>1</sub>	185	пър	Cys	, 41	001	190	1100	11011
			Asp		Glv	Va 1	Ser	Ala		Tle	Thr	Ile	Asn		Leu	Thr
254	0-1		195		0-1			200	E				205			
	Glv	Ser		Lvs	Thr	Pro	Glv		Ser	Ser	Val	Glv		Leu	Asp	Leu
258	U-1	210		-1-			215	1				220			<u>-</u>	
-	Gly	Gly	Gly	Ser	Thr	Gln	Ile	Ala	Phe	Leu	Pro	Arg	Val	Glu	Gly	Thr
	225	-	-			230					235	_			-	240
265	Leu	Gln	Ala	Ser	Pro	Pro	Gly	Tyr	Leu	Thr	Ala	Leu	Arg	Met	Phe	Asn
266					245		-	-		250					255	
269	Arg	Thr	Tyr	Lys	Leu	Tyr	Ser	Tyr	Ser	Tyr	Leu	Gly	Leu	Gly	Leu	Met
270				260					265					270		
273	Ser	Ala	Arg	Leu	Ala	Ile	Leu	Gly	Gly	Val	Glu	Gly	Gln	${\tt Pro}$	Ala	Lys
274			275					280					285			
277	Asp	Gly	Lys	Glu	Leu	Val	Ser	Pro	Cys	Leu	Ser	Pro	Ser	Phe	Lys	Gly
278		290					295					300				
281	Glu	Trp	Glu	His	Ala	Glu	Val	Thr	Tyr	Arg	Val	Ser	Gly	Gln	Lys	Ala
	305					310					315					320
	Ala	Ala	Ser	Leu	His	Glu	Leu	Cys	Ala		Arg	Val	Ser	Glu	Val	Leu
286					325					330					335	
	Gln	Asn	Arg		His	Arg	Thr	Glu		Val	Lys	His	Val		Phe	Tyr
290				340					345					350		
	Ala	Phe		$\mathtt{Tyr}$	$\mathtt{Tyr}$	Tyr	Asp		Ala	Ala	Gly	Val	_	Leu	Ile	Asp
294			355			_	_	360			_		365			_ •
	Ala		Lys	GŢĀ	СТА	Ser		Val	Val	GTA	Asp		Glu	IIe	Ala	Ala
298	_	370		_	_	_,	375	~ 7	<b>-1</b>		_	380		_	_	-1
	_	Tyr	Val	Cys	Arg		Leu	GLu	Thr	GIn		Gin	Ser	Ser	Pro	
	385	<b>a</b> -			<b>.</b>	390	m- ·	77- 3	a	+	395	<b>.</b>	<b>a</b> 7 ·	<b>a</b> 2	D1	400
	ser	Cys	мet	Asp		rnr	Tyr	vaı	ser		ьeu	ьeu	GIn	GIU	Phe	GTÀ
306	n1	D	3	0	405	17- 1	T	T	T	410	<b>3</b>	T	<b>-</b> 7 -	3	415	17- 7
309	Pne	Pro	Arg	ser	ьys	val	ьeu	гàг	ьeu	Thr	Arg	ьys	тте	Asp	Asn	val

**RAW SEQUENCE LISTING**PATENT APPLICATION: US/09/905,743B

DATE: 04/21/2003

TIME: 13:14:11

Input Set : N:\EBONY'S\EP.txt

Output Set: N:\CRF4\04212003\I905743B.raw

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     317 Asn Arg Gln Lys Ser Pro Ala Ser
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     321 <210> SEQ ID NO: 3
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     323 <212> TYPE: DNA
     324 <213> ORGANISM: Homo Sapiens
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     327 <221> NAME/KEY: CDS
     328 <222> LOCATION: (83)..(1669)
     329 <223> OTHER INFORMATION:
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     333 acceaegeqt ctggccqcgq geegectetg eggeageget agtegeette teegaategg
     335 ctccqcacag ctaggagaaa ag atg ttc act gtg ctg acc cgc caa cca tgt
                                                                                112
                                  Met Phe Thr Val Leu Thr Arg Gln Pro Cys
     336
     337
     339 gag caa gca ggc ctc aag gcc ctc tac cga act cca acc atc att gcc
                                                                                160
     340 Glu Gln Ala Gly Leu Lys Ala Leu Tyr Arg Thr Pro Thr Ile Ile Ala
     341
                         15
                                             20
                                                                                208
     343 ttq qtq qtc ttq ctt qtq aqt att qtq qta ctt qtq aqt atc act gtc
     344 Leu Val Val Leu Leu Val Ser Ile Val Val Leu Val Ser Ile Thr Val
                                         35
     347 atc cag atc cac aag caa gag gtc ctc cct cca gga ctg aag tat ggt
                                                                                256
     348 Ile Gln Ile His Lys Gln Glu Val Leu Pro Pro Gly Leu Lys Tyr Gly
                 45
                                     50
                                                                                304
     351 att gtg ctg gat gcc ggg tct tca aga acc aca gtc tac gtg tat caa
     352 Ile Val Leu Asp Ala Gly Ser Ser Arg Thr Thr Val Tyr Val Tyr Gln
     355 tgg cca gca gaa aaa gag aat aat acc gga gtg gtc agt caa acc ttc
                                                                                352
     356 Trp Pro Ala Glu Lys Glu Asn Asn Thr Gly Val Val Ser Gln Thr Phe
     359 aaa tgt agt gtg aaa ggc tct gga atc tcc agc tat gga aat aac ccc
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     360 Lys Cys Ser Val Lys Gly Ser Gly Ile Ser Ser Tyr Gly Asn Asn Pro
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                                             100
     363 caa gat gtc ccc aga gcc ttt gag gag tgt atg caa aaa gtc aag ggg
                                                                                448
     364 Gln Asp Val Pro Arg Ala Phe Glu Glu Cys Met Gln Lys Val Lys Gly
     365
                     110
                                         115
     367 cag gtt eca tee cae ete cae gga tee ace eee att cae etg gga gee
                                                                                496
     368 Gln Val Pro Ser His Leu His Gly Ser Thr Pro Ile His Leu Gly Ala
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                                     130
     371 acg gct ggg atg cgc ttg ctg agg ttg caa aat gaa aca gca gct aat
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     372 Thr Ala Gly Met Arg Leu Leu Arg Leu Gln Asn Glu Thr Ala Ala Asn
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                                 145
     375 gaa gtc ctt gaa agc atc caa agc tac ttc aag tcc cag ccc ttt gac
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     376 Glu Val Leu Glu Ser Ile Gln Ser Tyr Phe Lys Ser Gln Pro Phe Asp
     379 ttt agg ggt gct caa atc att tct ggg caa gaa gaa ggg gta tat gga
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**VERIFICATION SUMMARY** 

DATE: 04/21/2003 PATENT APPLICATION: US/09/905,743B TIME: 13:14:12

Input Set : N:\EBONY'S\EP.txt

Output Set: N:\CRF4\04212003\I905743B.raw

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